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Output Set: N:\CRF4\09122002\I782974C.raw

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132 Ile His Lys Asn Asp Gly Val Ser Leu Cys Phe Thr Leu Asn Leu Ala
136 Val Ala Asp Thr Leu Ile Gly Val Ala Ile Ser Gly Leu Leu Thr Asp
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| | 145 | | | | 150 | | | | | 155 | | _ | | | 160 | |
| 164 165 | Phe His | Pro | His | Phe 165 | Val | Leu | Thr | Leu | Ser 170 | Cys | Val | Gly | Phe | Phe 175 | Pro | |
| 168 169 | Ala Met | Leu | Leu 180 | Phe | Val | Phe | Phe | Tyr 185 | Cys | Asp | Met | Leu | Lys 190 | Ile | Ala | |
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| 173 | | 195 | | | | | 200 | | | | | 205 | _ | | _ | |
| 176 177 | Ala Gly 210 | _ | Tyr | Arg | Ser | Pro 215 | Arg | Thr | Pro | Ser | 220 | Phe | Lys | Ala | Leu | |
| 180 | Arg Thr | Val | Ser | Val | Ļeu | Ile | Gly | Ser | Phe | Ala | Leu | Ser | Trp | Thr | Pro | |
| 181 | 225 | | | | 230 | | | | | 235 | | | | | 240 | |
| 184 | Phe Leu | Ile | Thr | Gly | Ile | Val | Gln | Val | Ala | Cys | Gln | Glu | Cys | His | Leu | |
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| 189 | | | 260 | | | | | 265 | | | | | 270 | | | |
| 192 | Leu Leu | Asn | Pro | Leu | Ile | Tyr | Ala | Tyr | Trp | Gln | Lys | Glu | Val | Arg | Leu | |
| 193 | | 275 | | | | | 280 | | | | | 285 | | | | |
| 196 | Gln Leu | Tyr | His | Met | Ala | Leu | Gly | Val | Lys | Lys | Val | Leu | Thr | Ser | Phe | |
| 197 | 290 | | | | | 295 | | | | | 300 | | | | | |
| 200 | Leu Leu | Phe | Leu | Ser | Ala | Arg | Asn | Cys | Gly | Pro | Glu | Arg | Pro | Arg | | |
| | 305 | | | | 310 | | | | | 315 | | | | | 320 | |
| 204 | Ser Ser | Cys | His | Ile | Val | Thr | Ile | Ser | Ser | Ser | Glu | Phe | Asp | | | |
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| | gagctca | | | | | | | | | | | | | | | 600 |
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288 Thr Leu His Ala Val Gly Phe Val Leu Pro Leu Ala Val Leu Cys Leu
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| 378 Leu Pro Val Val Tyr Ser Leu Val Ala Ala Val Ser Ile Pro Gly Asn 379 | | | Leu Gln Met | | | | | | | | | | |
| 379 | | | • | | | | | | | | | | |
| 382 Leu Phe Ser Leu Trp Val Leu Cys Arg Arg Met Gly Pro Arg Ser Pro 383 | | | Tyr Ser Leu | | | | | | | | | | |
| 383 | | | _ | - - | | | | | | | | | |
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| 387 | | | | | * | | | | | | | | |
| 390 Ser Val Leu Pro Phe Gln Ile Tyr Tyr His Cys Asn Arg His His Trp 391 65 70 75 80 394 Val Phe Gly Val Leu Cys Asn Leu Val Val Thr Val Ala Phe Tyr Ala 395 85 90 95 398 Asn Met Tyr Ser Ser Ile Leu Thr Met Thr Cys Ile Ser Val Glu Arg 399 100 105 110 402 Phe Leu Gly Ile Leu Tyr Pro Leu Ser Ser Lys Arg Trp Arg Arg Arg 403 115 120 125 406 Arg Tyr Ala Val Ala Ala Cys Ala Gly Thr Trp Leu Leu Leu Thr 407 130 135 140 410 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala 411 145 150 155 160 414 Leu Gly Ile Ile Thr Cys Phe Asp Val 415 165 418 <210> SEQ ID No: 9 419 <211> LENGTH: 270 420 <212> TYPE: DNA 421 <213> ORGANISM: Homo sapiens 423 <400> SEQUENCE: 9 424 cccatgttcc tgctcctggg cagcctcacg ttgtcggatc tgctggagg cgccgctac 60 426 gccgccaca tcctactgtc ggggccgctc acgctgaaac tgctccccg gctctggttc 120 428 gcacgggagg gaggcgtctt cgtggcactc acgctgaaac tgctccccg gctctggttc 180 | | | | Leu Ser Val Thr | | | | | | | | | |
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| 395 | | | | | | | | | | | | | |
| 398 Asn Met Tyr Ser Ser Ile Leu Thr Met Thr Cys Ile Ser Val Glu Arg 399 100 105 110 402 Phe Leu Gly Ile Leu Tyr Pro Leu Ser Ser Lys Arg Trp Arg Arg Arg 403 115 120 125 406 Arg Tyr Ala Val Ala Ala Cys Ala Gly Thr Trp Leu Leu Leu Leu Thr 407 130 135 140 410 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala 411 145 150 155 160 414 Leu Gly Ile Ile Thr Cys Phe Asp Val 415 165 418 <210> SEQ ID NO: 9 419 <211> LENGTH: 270 420 <212> TYPE: DNA 421 <213> ORGANISM: Homo sapiens 423 <400> SEQUENCE: 9 424 cccatgttce tgctcctggg cagcctcacg ttgtcggatc tgctggcagg cgccgcctac 426 gccgccaaca tcctactgtc ggggccgctc acgcgctac tgctggcagg gaggcgtctt cgtggcact acgctgaacc tgctggcact cctgggcatc 428 gcacgggagg gaggcgtctt cgtggcactc actgcgtccg tgctgagcct cctgggcatc | | Val Phe Gly Val | = | | | | | | | | | | |
| 399 100 105 110 402 Phe Leu Gly Ile Leu Tyr Pro Leu Ser Ser Lys Arg Trp Arg Arg Arg 120 125 406 Arg Tyr Ala Val Ala Ala Cys Ala Gly Thr Trp Leu Leu Leu Leu Thr 130 135 140 410 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala 111 145 150 155 160 414 Leu Gly Ile Ile Thr Cys Phe Asp Val 165 165 165 165 168 168 169 169 169 169 160 1 | | | | | | | | | | | | | |
| 402 Phe Leu Gly Tle Leu Tyr Pro Leu Ser Ser Lys Arg Trp Arg Arg Arg 403 115 120 125 406 Arg Tyr Ala Val Ala Ala Cys Ala Gly Thr Trp Leu Leu Leu Leu Thr 407 130 135 140 410 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala 411 145 150 155 160 414 Leu Gly Ile Ile Thr Cys Phe Asp Val 415 165 418 <210> SEQ ID NO: 9 419 <211> LENGTH: 270 420 <212> TYPE: DNA 421 <213> ORGANISM: Homo sapiens 423 <400> SEQUENCE: 9 424 cccatgttcc tgctcctggg cagcctcacg ttgtcggatc tgctggagg cgccgctac 426 gccgccaaca tcctactgtc ggggccgctc acgctgaaac tgctccccgc gctctggttc 120 428 gcacgggagg gaggcgtctt cgtggcactc actgcgtccg tgctgagcct cctgggcatc 180 | | | ser lie Leu | | | | | | | | | | |
| 403 | | | Tou Mir Dro | , | | | | | | | | | |
| 406 Arg Tyr Ala Val Ala Ala Cys Ala Gly Thr Trp Leu Leu Leu Leu Thr 407 130 135 140 410 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala 411 145 150 155 160 414 Leu Gly Ile Ile Thr Cys Phe Asp Val 415 165 418 <210> SEQ ID NO: 9 419 <211> LENGTH: 270 420 <212> TYPE: DNA 421 <213> ORGANISM: Homo sapiens 423 <400> SEQUENCE: 9 424 cccatgttce tgctcctggg cagcctcacg ttgtcggatc tgctggagg cgccgctac 426 gccgccaaca tcctactgtc ggggccgctc acgctgaaac tgtcccccgc gctctggttc 428 gcacgggagg gaggcgtctt cgtggcactc actgcgtccg tgctgagcct cctgggcatc 180 | | _ | Led Tyl Plo | | | | | | | | | | |
| 407 130 135 140 410 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala 411 145 150 155 160 414 Leu Gly Ile Ile Thr Cys Phe Asp Val 415 165 418 <210> SEQ ID NO: 9 419 <211> LENGTH: 270 420 <212> TYPE: DNA 421 <213> ORGANISM: Homo sapiens 423 <400> SEQUENCE: 9 424 cccatgttcc tgctcctggg cagcctcacg ttgtcggatc tgctggcagg cgccgctac 426 gccgccaaca tcctactgtc ggggccgctc acgctgaacc tgctcccgc gctctggttc 428 gcacgggagg gaggcgtctt cgtggcactc actgcgccg tgctgagcct cctgggcatc 180 | | | Ala Ala Crra | | | | | | | | | | |
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| 424 cccatgttcc tgctcctggg cagcctcacg ttgtcggatc tgctggcagg cgccgctac60426 gccgccaaca tcctactgtc ggggccgctc acgctgaaac tgtcccccgc gctctggttc120428 gcacgggagg gaggcgtctt cgtggcactc actgcgtccg tgctgagcct cctgggcatc180 | | | | | | | | | | | | | |
| 426 gccgccaaca tcctactgtc ggggccgctc acgctgaaac tgtcccccgc gctctggttc 120 428 gcacgggagg gaggcgtctt cgtggcactc actgcgtccg tgctgagcct cctgggcatc 180 | | | | | | | | | | | | | |
| 428 gcacgggagg gaggcgtctt cgtggcactc actgcgtccg tgctgagcct cctgggcatc 180 | | | | | | | | | | | | | |
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/782,974C

DATE: 09/12/2002 TIME: 15:27:59

Input Set : A:\00411PHRM311.ST25.txt Output Set: N:\CRF4\09122002\I782974C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:15; N Pos. 431,432,433,434,435,436,437,438,439,440,441,442,443,444,445 Seq#:15; N Pos. 446,447,448,449,450,451,452,453,454,455,456,457,458,459,460 Seq#:15; N Pos. 461,790,791,792,793,794,795,796,797,798,799,800,801,802,803 Seq#:15; N Pos. 804,805,806,807,808,809,810,811,812,813,814,815,816,817,818

Seq#:16; Xaa Pos. 26,27,28,29,30,31,32,33,34,35,144,145,146,147,148,149,150

Seq#:16; Xaa Pos. 151,152,153,154

 $\texttt{Seq} \# : 27; \ \texttt{N} \ \texttt{Pos.} \ 81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99$

Seq#:27; N Pos. 100,101,102,103,104,105,106

Seq#:28; Xaa Pos. 104,105,106,107,108,109,110,111,112,113

Seq#:58; Xaa Pos. 266